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Ile Phe Lys Gln Ile Ala Gly Asp Asp Met Glu Ile Cys Ala Asp Glu
            660              665              670

Leu Lys Lys Val Leu Asn Thr Val Val Asn Lys His Lys Asp Leu Lys
            675              680              685

Thr His Gly Phe Thr Leu Glu Ser Cys Arg Ser Met Ile Ala Leu Met
            690              695              700

Asp Thr Asp Gly Ser Gly Lys Leu Asn Leu Gln Glu Phe His His Leu
705              710              715              720

Trp Asn Lys Ile Lys Ala Trp Gln Lys Ile Phe Lys His Tyr Asp Thr

                              20
```

```
              725                    730                    735

Asp Gln Ser Gly Thr Ile Asn Ser Tyr Glu Met Arg Asn Ala Val Asn
            740                    745                    750

Asp Ala Gly Phe His Leu Asn Asn Gln Leu Tyr Asp Ile Ile Thr Met
            755                    760                    765

Arg Tyr Ala Asp Lys His Met Asn Ile Asp Phe Asp Ser Phe Ile Cys
            770                    775                    780

Cys Phe Val Arg Leu Glu Gly Met Phe Arg Ala Phe His Ala Phe Asp
785                    790                    795                    800

Lys Asp Gly Asp Gly Ile Ile Lys Leu Asn Val Leu Glu Trp Leu Gln
                   805                    810                    815

Leu Thr Met Tyr Ala
                820


<210> 7
<211> 821
<212> PRT
<213> Homo sapiens

<220>
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    /note= "FIGURE 3 - RAT"

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21

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/note= "FIGURE 3 - RAT"

<400> 7

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Met Pro Thr Val Ile Ser Pro Thr Val Ala Pro Arg Thr Gly Ala Glu
 1               5                  10                  15

Pro Arg Ser Pro Gly Pro Val Pro His Pro Ala Gln Gly Lys Thr Thr
                20                  25                  30

Glu Ala Gly Gly Gly His Pro Gly Gly Ile Tyr Ser Ala Ile Ile Ser
            35                  40                  45

Arg Asn Phe Pro Ile Ile Gly Val Lys Glu Lys Thr Phe Glu Gln Leu
        50                  55                  60

His Lys Lys Cys Leu Glu Lys Lys Val Leu Tyr Leu Asp Pro Glu Phe
 65                  70                  75                  80

Pro Pro Asp Glu Thr Ser Leu Phe Tyr Ser Gln Lys Phe Pro Ile Gln
                85                  90                  95

Phe Val Trp Lys Arg Pro Pro Glu Ile Cys Glu Asn Pro Arg Phe Ile
               100                 105                 110

Ile Gly Gly Ala Asn Arg Thr Asp Ile Cys Gln Gly Asp Leu Gly Asp
           115                 120                 125

Cys Trp Leu Leu Ala Ala Ile Ala Cys Leu Thr Leu Asn Glu Arg Leu
       130                 135                 140

Leu Phe Arg Val Ile Pro His Asp Gln Ser Phe Thr Glu Asn Tyr Ala
145                 150                 155                 160

Gly Ile Phe His Phe Gln Phe Trp Arg Tyr Gly Asp Trp Val Asp Val
               165                 170                 175
```

22

Val Ile Asp Asp Cys Leu Pro Thr Tyr Asn Asn Gln Leu Val Phe Thr
            180   185   190

Lys Ser Asn His Arg Asn Glu Phe Trp Ser Ala Leu Leu Glu Lys Ala
            195   200   205

Tyr Ala Lys Leu His Gly Ser Tyr Glu Ala Leu Lys Gly Gly Asn Thr
    210   215   220

Thr Glu Ala Met Glu Asp Phe Thr Gly Gly Val Thr Glu Phe Phe Glu
225   230   235   240

Ile Lys Asp Ala Pro Ser Asp Met Tyr Lys Ile Met Arg Lys Ala Ile
                245   250   255

Glu Arg Gly Ser Leu Met Gly Cys Ser Ile Asp Asp Gly Thr Asn Met
            260   265   270

Thr Tyr Gly Thr Ser Pro Ser Gly Leu Asn Met Gly Glu Leu Ile Ala
            275   280   285

Arg Met Val Arg Asn Met Asp Asn Ser Leu Leu Arg Asp Ser Asp Leu
    290   295   300

Asp Pro Arg Ala Ser Asp Asp Arg Pro Ser Arg Thr Ile Val Pro Val
305   310   315   320

Gln Tyr Glu Thr Arg Met Ala Cys Gly Leu Val Arg Gly His Ala Tyr
                325   330   335

Ser Val Thr Gly Leu Glu Glu Ala Leu Phe Lys Gly Glu Lys Val Lys
            340   345   350

Leu Val Arg Leu Arg Asn Pro Trp Gly Gln Val Glu Trp Asn Gly Ser
            355   360   365

Trp Ser Asp Gly Trp Lys Asp Trp Ser Phe Val Asp Lys Asp Glu Lys
    370   375   380

Ala Arg Leu Gln His Gln Val Thr Glu Asp Gly Glu Phe Trp Met Ser
385   390   395   400

Tyr Asp Asp Phe Val Tyr His Phe Thr Lys Leu Glu Ile Cys Asn Leu
                405   410   415

Thr Ala Asp Ala Leu Glu Ser Asp Lys Leu Gln Thr Trp Thr Val Ser
            420   425   430

23

```
Val Asn Glu Gly Arg Trp Val Arg Gly Cys Ser Ala Gly Gly Cys Arg
        435                 440             445

Asn Phe Pro Asp Thr Phe Trp Thr Asn Pro Gln Tyr Arg Leu Lys Leu
        450             455             460

Leu Glu Glu Asp Asp Asp Pro Asp Asp Ser Glu Val Ile Cys Ser Phe
465             470             475                 480

Leu Val Ala Leu Met Gln Lys Asn Arg Arg Lys Asp Arg Lys Leu Gly
            485             490             495

Ala Asn Leu Phe Thr Ile Gly Phe Ala Ile Tyr Glu Val Pro Lys Glu
            500             505             510

Met His Gly Asn Lys Gln His Leu Gln Lys Asp Phe Phe Leu Tyr Asn
        515             520             525

Ala Ser Lys Ala Arg Ser Lys Thr Tyr Ile Asn Met Arg Glu Val Ser
        530             535             540

Gln Arg Phe Arg Leu Pro Pro Ser Glu Tyr Val Ile Val Pro Ser Thr
545             550             555             560

Tyr Glu Pro His Gln Glu Gly Glu Phe Ile Leu Arg Val Phe Ser Glu
            565             570             575

Lys Arg Asn Leu Ser Glu Glu Ala Glu Asn Thr Ile Ser Val Asp Arg
            580             585             590

Pro Val Lys Lys Lys Lys Asn Lys Pro Ile Ile Phe Val Ser Asp Arg
            595             600             605

Ala Asn Ser Asn Lys Glu Leu Gly Val Asp Gln Glu Ala Glu Glu Gly
        610             615             620

Lys Asp Lys Thr Gly Pro Asp Lys Gln Gly Glu Ser Pro Gln Pro Arg
625             630             635             640

Pro Gly His Thr Asp Gln Glu Ser Glu Glu Gln Gln Gln Phe Arg Asn
            645             650             655

Ile Phe Arg Gln Ile Ala Gly Asp Asp Met Glu Ile Cys Ala Asp Glu
            660             665             670

Leu Lys Asn Val Leu Asn Thr Val Val Asn Lys His Lys Asp Leu Lys
        675             680             685
```

24

Thr Gln Gly Phe Thr Leu Glu Ser Cys Arg Ser Met Ile Ala Leu Met
    690   695   700

Asp Thr Asp Gly Ser Gly Arg Leu Asn Leu Gln Glu Phe His His Leu
705   710   715   720

Trp Lys Lys Ile Lys Ala Trp Gln Lys Ile Phe Lys His Tyr Asp Thr
            725   730   735

Asp His Ser Gly Thr Ile Asn Ser Tyr Glu Met Arg Asn Ala Val Asn
            740   745   750

Asp Ala Gly Phe His Leu Asn Ser Gln Leu Tyr Asp Ile Ile Thr Met
            755   760   765

Arg Tyr Ala Asp Lys His Met Asn Ile Asp Phe Asp Ser Phe Ile Cys
    770   775   780

Cys Phe Val Arg Leu Glu Gly Met Phe Arg Ala Phe His Ala Phe Asp
785   790   795   800

Lys Asp Gly Asp Gly Ile Ile Lys Leu Asn Val Leu Glu Trp Leu Gln
            805   810   815

Leu Thr Met Tyr Ala
            820

<210> 8
<211> 821
<212> PRT
<213> Homo sapiens

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    /note= "FIGURE 3 -PIG"

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/note= "FIGURE 3 -PIG"

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/note= "FIGURE 3 -PIG"

<400> 8
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Met Pro Thr Val Ile Ser Ala Ser Val Ala Pro Arg Thr Ala Ala Glu
  1               5               10              15

Pro Arg Ser Pro Gly Pro Val Pro His Pro Ala Gln Ser Lys Ala Thr
             20              25              30

Glu Ala Gly Gly Gly Asn Pro Ser Gly Ile Tyr Ser Ala Ile Ile Ser
         35              40              45

Arg Asn Phe Pro Ile Ile Gly Val Lys Glu Lys Thr Phe Glu Gln Leu
     50              55              60

His Lys Lys Cys Leu Glu Lys Lys Val Leu Tyr Val Asp Pro Glu Phe
 65              70              75              80

Pro Pro Asp Glu Thr Ser Leu Phe Tyr Ser Gln Lys Phe Pro Ile Gln
             85              90              95

Phe Val Trp Lys Arg Pro Pro Glu Ile Cys Glu Asn Pro Arg Phe Ile
            100             105             110
```

26

```
Ile Asp Gly Ala Asn Arg Thr Asp Ile Cys Gln Gly Glu Leu Gly Asp
        115                 120                 125

Cys Trp Phe Leu Ala Ala Ile Ala Cys Leu Thr Leu Asn Gln His Leu
        130                 135                 140

Leu Phe Arg Val Ile Pro His Asp Gln Ser Phe Ile Glu Asn Tyr Ala
145                 150                 155                 160

Gly Ile Phe His Phe Gln Phe Trp Arg Tyr Gly Glu Trp Val Asp Val
                165                 170                 175

Val Ile Asp Asp Cys Leu Pro Thr Tyr Asn Asn Gln Leu Val Phe Thr
                180                 185                 190

Lys Ser Asn His Arg Asn Glu Phe Trp Ser Ala Leu Leu Glu Lys Ala
        195                 200                 205

Tyr Ala Lys Leu His Gly Ser Tyr Glu Ala Leu Lys Gly Gly Asn Thr
    210                 215                 220

Thr Glu Ala Met Glu Asp Phe Thr Gly Gly Val Ala Glu Phe Phe Glu
225                 230                 235                 240

Ile Arg Asp Ala Pro Ser Asp Met Tyr Lys Ile Met Lys Lys Ala Ile
                245                 250                 255

Glu Arg Gly Ser Leu Met Gly Cys Ser Ile Asp Asp Gly Thr Asn Met
                260                 265                 270

Thr Tyr Gly Thr Ser Pro Ser Gly Leu Asn Met Gly Asp Leu Ile Ala
        275                 280                 285

Arg Met Val Arg Asn Met Glu Asn Ser Arg Leu Arg Asp Ser Ile Leu
        290                 295                 300

Asp Pro Glu Val Ser Asp Asp Arg Pro Thr Arg Thr Ile Val Pro Val
305                 310                 315                 320

Gln Phe Glu Thr Arg Met Ala Cys Gly Leu Val Arg Gly His Ala Tyr
                325                 330                 335

Ser Val Thr Gly Leu Glu Glu Ala Leu Phe Lys Gly Glu Lys Val Lys
            340                 345                 350

Leu Val Arg Leu Arg Asn Pro Trp Gly Gln Val Glu Trp Asn Gly Ser
            355                 360                 365

                              27
```

Trp Ser Asp Ser Trp Lys Asp Trp Ser Phe Val Asp Lys Asp Glu Lys
    370   375   380

Ala Arg Leu Gln His Gln Val Thr Glu Asp Gly Glu Phe Trp Met Ser
385   390   395   400

Tyr Asp Asp Phe Ile Tyr His Phe Thr Lys Leu Glu Ile Cys Asn Leu
            405   410   415

Thr Ala Asp Ala Leu Glu Ser Asp Lys Leu Gln Thr Trp Thr Val Ser
            420   425   430

Val Asn Glu Gly Arg Trp Val Arg Gly Cys Ser Ala Gly Thr Gly Arg
        435   440   445

Asn Phe Pro Asp Thr Phe Trp Thr Asn Pro Gln Tyr Arg Leu Lys Leu
    450   455   460

Leu Glu Glu Asp Asp Asp Pro Asp Asp Ser Glu Val Ile Cys Ser Phe
465   470   475   480

Leu Val Ala Leu Met Gln Arg Asn Arg Arg Lys Asp Arg Lys Leu Gly
            485   490   495

Ala Asn Leu Phe Thr Ile Gly Phe Ala Ile Tyr Glu Val Pro Lys Glu
        500   505   510

Met His Gly Asn Lys Gln His Leu Gln Lys Asp Phe Phe Leu Tyr Asn
        515   520   525

Ala Ser Lys Ala Arg Ser Arg Thr Tyr Ile Asn Met Arg Glu Val Ser
    530   535   540

Glu Arg Phe Arg Leu Pro Pro Ser Glu Tyr Val Ile Val Pro Ser Thr
545   550   555   560

Tyr Glu Pro His Gln Glu Gly Glu Phe Met Leu Arg Val Phe Ser Glu
            565   570   575

Lys Arg Lys Leu Ser Glu Glu Val Glu Asn Thr Ile Ser Val Asp Arg
        580   585   590

Pro Val Arg Lys Lys Lys Thr Lys Pro Ile Ile Phe Val Ser Asp Arg
        595   600   605

Ala Asn Ser Asn Lys Glu Leu Gly Val Asp Gln Glu Ser Glu Glu Gly
    610   615   620

28

```
Gln Asp Lys Thr Ser Pro Asp Lys Gln Glu Lys Ser Pro Lys Pro Glu
625                 630                 635                 640

Pro Ser Asn Thr Asp Gln Glu Ser Glu Glu Gln Gln Gln Phe Arg Asn
                645                 650                 655

Ile Phe Lys Gln Ile Ala Gly Asp Asp Met Glu Ile Cys Ala Asp Glu
            660                 665                 670

Leu Lys Lys Val Leu Asn Thr Val Val Asn Lys His Lys Asp Leu Lys
        675                 680                 685

Thr His Gly Phe Thr Leu Glu Ser Cys Arg Ser Met Ile Ala Leu Met
    690                 695                 700

Asp Thr Asp Gly Ser Gly Lys Leu Asn Leu Gln Glu Phe His His Leu
705                 710                 715                 720

Trp Asn Lys Ile Lys Ala Trp Gln Lys Ile Phe Lys His Tyr Asp Thr
            725                 730                 735

Asp Gln Ser Gly Thr Ile Asn Ser Tyr Glu Met Arg Asn Ala Val Asn
            740                 745                 750

Asp Ala Gly Phe His Leu Asn Asn Gln Leu Tyr Asp Ile Ile Thr Met
            755                 760                 765

Arg Tyr Ala Asp Lys His Met Asn Ile Asp Phe Asp Ser Phe Ile Cys
    770                 775                 780

Cys Phe Val Arg Leu Glu Gly Met Phe Arg Ala Phe His Ala Phe Asp
785                 790                 795                 800

Lys Asp Gly Asp Gly Ile Ile Lys Leu Asn Val Leu Glu Trp Leu Gln
            805                 810                 815

Leu Thr Met Tyr Ala
            820

<210> 9
<211> 821
<212> PRT
<213> Homo sapiens

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/note= "FIGURE 3 -COW"

<220>
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/note= "FIGURE 3 -COW"

<220>
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/note= "FIGURE 3 -COW"

<400> 9
Met Pro Thr Val Ile Ser Ala Ser Val Ala Pro Arg Thr Ala Ala Glu
1   5   10   15

Pro Arg Ser Pro Gly Pro Val Pro His Pro Ala Gln Ser Lys Ala Thr
                20   25   30

Glu Ala Gly Gly Gly Asn Pro Ser Gly Ile Tyr Ser Ala Ile Ile Ser
            35   40   45

Arg Asn Phe Pro Ile Ile Gly Val Lys Glu Lys Thr Phe Glu Gln Leu

```
          50                    55                    60

His Lys Lys Cys Leu Glu Lys Lys Val Leu Tyr Val Asp Pro Glu Phe
65              70              75                        80

Pro Pro Asp Glu Thr Ser Leu Phe Tyr Ser Gln Lys Phe Pro Ile Gln
                85              90                        95

Phe Val Trp Lys Arg Pro Pro Glu Ile Cys Glu Asn Pro Arg Phe Ile
            100             105             110

Ile Asp Gly Ala Asn Arg Thr Asp Ile Cys Gln Gly Glu Leu Gly Asp
        115             120             125

Cys Trp Phe Leu Ala Ala Ile Ala Cys Leu Thr Leu Asn Gln His Leu
    130             135             140

Leu Phe Arg Val Ile Pro His Asp Gln Ser Phe Ile Glu Asn Tyr Ala
145             150             155                       160

Gly Ile Phe His Phe Gln Phe Trp Arg Tyr Gly Glu Trp Val Asp Val
            165             170             175

Val Ile Asp Asp Cys Leu Pro Thr Tyr Asn Asn Gln Leu Val Phe Thr
            180             185             190

Lys Ser Asn His Arg Asn Glu Phe Trp Ser Ala Leu Leu Glu Lys Ala
        195             200             205

Tyr Ala Lys Leu His Gly Ser Tyr Glu Ala Leu Lys Gly Gly Asn Thr
    210             215             220

Thr Glu Ala Met Glu Asp Phe Thr Gly Gly Val Ala Glu Phe Phe Glu
225             230             235             240

Ile Arg Asp Ala Pro Ser Asp Met Tyr Lys Ile Met Lys Lys Ala Ile
            245             250             255

Glu Arg Gly Ser Leu Met Gly Cys Ser Ile Asp Asp Gly Thr Asn Met
        260             265             270

Thr Tyr Gly Thr Ser Pro Ser Gly Leu Asn Met Gly Glu Leu Ile Glu
        275             280             285

Arg Met Val Arg Asn Met Asp Asn Ser Arg Leu Arg Asp Ser Ile Leu
    290             295             300

Asp Pro Glu Val Ser Asp Asp Arg Pro Thr Arg Met Ile Val Pro Val

                                31
```

```
                    305                        310                         315                         320

Gln Phe Glu Thr Arg Met Ala Cys Gly Leu Val Arg Gly His Ala Tyr
              325                        330                        335

Ser Val Thr Gly Leu Glu Glu Ala Leu Tyr Lys Gly Glu Lys Val Lys
              340                        345                        350

Leu Val Arg Leu Arg Asn Pro Trp Gly Gln Val Glu Trp Asn Gly Ser
              355                        360                        365

Trp Ser Asp Ser Trp Lys Asp Trp Ser Tyr Val Asp Lys Asp Glu Lys
     370                        375                        380

Ala Arg Leu Gln His Gln Val Thr Glu Asp Gly Glu Phe Trp Met Ser
385                        390                        395                        400

Tyr Glu Asp Phe Ile Tyr His Phe Thr Lys Leu Glu Ile Cys Asn Leu
              405                        410                        415

Thr Ala Asp Ala Leu Gln Ser Asp Lys Leu Gln Thr Trp Thr Val Ser
              420                        425                        430

Val Asn Glu Gly Arg Trp Val Arg Gly Cys Ser Ala Gly Gly Cys Arg
              435                        440                        445

Asn Phe Pro Asp Thr Phe Trp Thr Asn Pro Gln Tyr Arg Leu Lys Leu
              450                        455                        460

Leu Glu Glu Asp Asp Asp Pro Asp Asp Ser Glu Val Ile Cys Ser Phe
465                        470                        475                        480

Leu Val Ala Leu Met Gln Lys Asn Arg Arg Lys Asp Arg Lys Leu Gly
              485                        490                        495

Ala Ser Leu Phe Thr Ile Gly Phe Ala Ile Tyr Glu Val Pro Lys Glu
              500                        505                        510

Met His Gly Asn Lys Gln His Leu Gln Lys Asp Phe Phe Leu Tyr Asn
              515                        520                        525

Ala Ser Lys Ala Arg Ser Lys Thr Tyr Ile Asn Met Arg Glu Val Ser
              530                        535                        540

Gln Arg Phe Arg Leu Pro Pro Ser Glu Tyr Val Ile Val Pro Ser Thr
545                        550                        555                        560

Tyr Glu Pro His Gln Glu Gly Glu Phe Ile Leu Arg Val Phe Ser Glu
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32

Lys Arg Asn Leu Ser Glu Glu Val Glu Asn Thr Ile Ser Val Asp Arg
            580   585   590

Pro Val Lys Lys Lys Lys Thr Lys Pro Ile Ile Phe Val Ser Asp Arg
            595   600   605

Ala Asn Ser Asn Lys Glu Leu Gly Val Asp Gln Glu Ser Glu Glu Gly
            610   615   620

Lys Gly Lys Thr Ser Pro Asp Lys Gln Lys Gln Ser Pro Gln Pro Gln
625   630   635   640

Pro Gly Ser Ser Asp Gln Glu Ser Glu Glu Gln Gln Gln Phe Arg Asn
                645   650   655

Ile Phe Lys Gln Ile Ala Gly Asp Asp Met Glu Ile Cys Ala Asp Glu
            660   665   670

Leu Lys Lys Val Leu Asn Thr Val Val Asn Lys His Lys Asp Leu Lys
            675   680   685

Thr His Gly Phe Thr Leu Glu Ser Cys Arg Ser Met Ile Ala Leu Met
            690   695   700

Asp Thr Asp Gly Ser Gly Lys Leu Asn Leu Gln Glu Phe His His Leu
705   710   715   720

Trp Asn Lys Ile Lys Ala Trp Gln Lys Ile Phe Lys His Tyr Asp Thr
                725   730   735

Asp Gln Ser Gly Thr Ile Asn Ser Tyr Glu Met Arg Asn Ala Val Asn
                740   745   750

Asp Ala Gly Phe His Leu Asn Asn Gln Leu Tyr Asp Ile Ile Thr Met
                755   760   765

Arg Tyr Ala Asp Lys His Met Asn Ile Asp Phe Asp Ser Phe Ile Cys
                770   775   780

Cys Phe Val Arg Leu Glu Gly Met Phe Arg Ala Phe His Ala Phe Asp
785   790   795   800

Lys Asp Gly Asp Gly Ile Ile Lys Leu Asn Val Leu Glu Trp Leu Gln
                805   810   815

Leu Thr Met Tyr Ala

33

<210> 10
<211> 20
<212> DNA
<213> Homo sapiens

<220>
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<210> 11
<211> 21
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(21)
<223> /label= Table 1

<400> 11
gtatgactcg gaaaagaagg t   21

<210> 12
<211> 20
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(20)
<223> /label= Table 1

<400> 12
taagcaaaag cagtccccac   20

<210> 13
<211> 21
<212> DNA
<213> Homo sapiens

<220>

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<221> misc_feature
<222> (1)..(21)
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<400> 13
ttgctgttcc tcactttcct g                                    21

<210> 14
<211> 20
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(20)
<223> /label= Table 1

<400> 14
gtttcatctg ctgcttcgtt                                      20

<210> 15
<211> 20
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(20)
<223> /label= Table 1

<400> 15
ctggttcagg catacatggt                                      20

<210> 16
<211> 22
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(22)
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<400> 16
ttctttatgt ggaccctgag tt                                   22

<210> 17
<211> 19
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35

<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(19)
<223> /label= Table 1

<400> 17
acgaactgga tggggaact   19

<210> 18
<211> 20
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(20)
<223> /label= Table 3

<400> 18
ttcagtacct cccgttcacc   20

<210> 19
<211> 20
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(20)
<223> /label= Table 3

<400> 19
gatgcttgag ccaggaaaac   20

<210> 20
<211> 22
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)..(22)
<223> /label= Table 3

<400> 20

36

ctttccttga aggtagctgt at   22

<210> 21
<211> 20
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)..(20)
<223> /label= Table 3

<400> 21
gaggtgctga gtgagaggac   20

<210> 22
<211> 22
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(22)
<223> /label= Table 3

<400> 22
actccgtctc aaaaaaatac ct   22

<210> 23
<211> 20
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(20)
<223> /label= Table 3

<400> 23
attgtccctt tacctcctgg   20

<210> 24
<211> 20
<212> DNA
<213> Homo sapiens

<220>

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<221> misc_feature
<222> (1)..(20)
<223> /label= Table 3


<400> 24
tggaagtagg agagtgggca                                          20


<210> 25
<211> 20
<212> DNA
<213> Homo sapiens


<220>
<221> misc_feature
<222> (1)..(20)
<223> /label= Table 3


<400> 25
gggtagatgg gtgggaagtt                                          20


<210> 26
<211> 20
<212> DNA
<213> Homo sapiens


<220>
<221> misc_feature
<222> (1)..(20)
<223> /label= Table 3


<400> 26
gaggaatgtg gaggaaggac                                          20


<210> 27
<211> 20
<212> DNA
<213> Homo sapiens


<220>
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<222> (1)..(20)
<223> /label= Table 3


<400> 27
ttcctgtgag tgaggtctcg                                          20


<210> 28
<211> 20
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38

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<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)..(20)
<223> /label= Table 3

<400> 28
ggaactctgt gaccccaaat                                              20

<210> 29
<211> 20
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)..(20)
<223> /label= Table 3

<400> 29
tcctcaaaca aaacattcgc                                              20

<210> 30
<211> 20
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(20)
<223> /label= Table 3

<400> 30
gttccctaca ttctccatcg                                              20

<210> 31
<211> 21
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(21)
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<400> 31
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Sub
D(
at
cont

gttatttcaa cccagaccct t   21

<210> 32
<211> 21
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(21)
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<400> 32
aatgggttct ctggttactg c   21

<210> 33
<211> 21
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)..(21)
<223> /label= Table 3

<400> 33
agcacgaaaa gcaaagataa a   21

<210> 34
<211> 20
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)..(20)
<223> /label= Table 3

<400> 34
gtaagagatt tgccccccag   20

<210> 35
<211> 20
<212> DNA
<213> Homo sapiens

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tgtggatttg tgtgctacgc   20

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